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Page 1 of 6

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1600

RAW SEQUENCE LISTING

DATE: 09/11/2002

PATENT APPLICATION: US/09/383,579B

TIME: 10:38:41

Input Set : A:\2531-1-001revised.ST25.txt

Output Set : N:\CRF4\09112002\I383579B.raw

3 <110> APPLICANT: Burrell, Michael M.
4 Cambridge, Amanda P.
5 Maunders, Martin J.
6 McQueen-Mason, Simon
8 <120> TITLE OF INVENTION: Modification of Plant Fibres
10 <130> FILE REFERENCE: 2531-1-001
12 <140> CURRENT APPLICATION NUMBER: 09/383,579B
13 <141> CURRENT FILING DATE: 1999-08-25
15 <150> PRIOR APPLICATION NUMBER: UK 9818808.9
16 <151> PRIOR FILING DATE: 1998-08-29
18 <160> NUMBER OF SEQ ID NOS: 8
20 <170> SOFTWARE: PatentIn version 3.1
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 488
24 <212> TYPE: DNA
25 <213> ORGANISM: eucalyptus grandis
27 <400> SEQUENCE: 1
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30 gctttgagca ctgccctgtt caacaatggc ctgagctgcg gggcatgtta cgagatgcgg 120
32 tgcaacgacg accccaggtg gtgcctcccg gggaccatca tggtcacggc aaccaacttt 180
34 tgccctccca acttggccct ctccaacgac aattgcggct ggtgcaaccc ccctctccag 240
36 cacttcgata tggccgagcc tgctttcttg cagattgccc agtaciaaagc tgggattgtc 300
38 caggtttccct tcagaagggt tccgtgtgtg aagaaaggag gggtaagggt caccatcaat 360
40 gggcactcct acttcaactt ggtgctgatc accaacgtgg gaggtgctgg tgatgtccat 420
42 tccgtttcca tcaagggctc gaggactggt tggcaagcca tgtcaaggaa ctggggcaaa 480
44 aactggca 488
47 <210> SEQ ID NO: 2
48 <211> LENGTH: 475
49 <212> TYPE: DNA
50 <213> ORGANISM: eucalyptus grandis
52 <400> SEQUENCE: 2
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55 gctttgagca ctgccctgtt caacaatggc ctgagctgcg gggcatgtta cgagatgcgg 120
57 tgcaacgacg accccaggtg gtgcctcccg gggaccatca tggtcacggc aaccaacttt 180
59 tgccctccca acttggccct ctccaacgac aatggcggct ggtgcaaccc ccctctccag 240
61 cacttcgata tggccgagcc tgctttcttg cagattgccc agtaciaaagc tgggattgtc 300
63 ccggtttcct tcagaagggt tccgtgtgtg aagaaaggag gggtaagggt caccatcaat 360
65 gggcactcct acttcagctg tgggtctgat caccaacgtg ggaggtgctg gtgatgtcca 420
67 ttccgtttcc atcaagagct cgaggactgg ttggcaagcc atgtcaaggaa attga 475
70 <210> SEQ ID NO: 3
71 <211> LENGTH: 494
72 <212> TYPE: DNA
73 <213> ORGANISM: eucalyptus grandis

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75 <400> SEQUENCE: 3
76 atggggggggg catgtggtta cggggacctt cacagggcca cctatggcaa gtacagtgcc      60
78 ggcttgagct cgatgctgtt caacagaggg agtacctgcg gggcttgctt cgagctccgg      120
80 tgcgtcgacc acattttgtg gtgcctccct ggtagcccggt cggatgcctt caccgccacc      180
82 gacttctgcc ctccgaacta cgggctcgcg gcagattacg gcgggtggtg caacttcccg      240
84 caggagcaact tcgagatgtc ggaggcgggc ttccgcccaga ttgcgggtgcg aagggtgat      300
86 gtggtgccta tccagtacag gagggtgaac tgtctgagaa gcgggtggtt gagattcaca      360
88 ttgagcggaa actctcactt ctttcaggtc ttggtgacga atgtaggcct agatggggag      420
90 gtgattgcca tgaaaatgaa gggatcgaaa acaggggtgga taccgatggc aagaaactgg      480
92 ggcaaaaact ggca
95 <210> SEQ ID NO: 4
96 <211> LENGTH: 437
97 <212> TYPE: DNA
98 <213> ORGANISM: eucalyptus grandis
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103 ccgacattgg ttatgagcac gaggttgaaa taagaatggc cgttgacggt gaaccggatc      120
105 cctccgcttc tctgcacct cactcttcgg taggccaccg ggacgatccc ggccctgtac      180
107 tgcgcaatgt gctggaagac cggctgggag aggtcgaaat ggagttgagg agggtcgcac      240
109 caccctcctg gagggcagaa gttggtcgcc gtgaccacaa tggcgcccgg gaggcaccac      300
111 tgcgggtcgt tcacgcaccg gagctcaaaag caccgcgcgc agctcagccc attggtgaac      360
113 aatgcagtgc tcagtgcagc tgtgtttgtg cgtaccctt ggctgtatag attcccataa      420
115 ccacacgccc ccccat
118 <210> SEQ ID NO: 5
119 <211> LENGTH: 437
120 <212> TYPE: DNA
121 <213> ORGANISM: eucalyptus grandis
123 <400> SEQUENCE: 5
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126 ccgacattgg ttatgagcac gaggttgaaa taagaatggc cgttgacggt gaaccggatc      120
128 cctccgcttc tctgcacct cactcttcgg taggccacag ggacgatccc ggccctgtac      180
130 tgcgcaatgt gctggaagac aggtcgaggag aggtcgaaat ggagttgagg agggtcgcac      240
132 caccctcctg gagggcagaa gttggtcgcc gtgacaacaa tggcgcccgg gaggcaccac      300
134 tgcgggtcgt tcacgcaccg gagctcaaaag caccgcgcgc agctcagccc attggtgaac      360
136 aatgcagtgc tcagtgcagc tgtgtttgtg cgtaccctt ggctgtatag attcccataa      420
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141 <210> SEQ ID NO: 6
142 <211> LENGTH: 448
143 <212> TYPE: DNA
144 <213> ORGANISM: eucalyptus grandis
146 <400> SEQUENCE: 6
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149 tgcgcgcgcg acgtttgtga cgaggacgag cttgaagtat gagttgccgt tgatggtgaa      120
151 ccggtatgct cctctcctcc tgcacgtcac cctcctgtac gcaacgtgga cgatgccggc      180
153 tcggtacttg gcaatgtgct ggaagacggg ctgggagatg tcgaagtggg gttggggcgg      240
155 gttgcaccat ccgcggcggt tgtttgggag ggcgttgttt ggcgggcaga agtttgtggc      300
157 ggtgacgacg atggagccgc ccaggcacca ctttcctcgc ttcacgcacc ggatctcgaa      360
159 gcacgacccc cagctcagcc cgttttttaa cagcgccgtg ctcagcgccg ccgtgttcgt      420
161 accgtagccc tggctgtaca ggttgccg

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TIME: 10:38:41

Input Set : A:\2531-1-001revised.ST25.txt

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164 <210> SEQ ID NO: 7
165 <211> LENGTH: 19
166 <212> TYPE: DNA
167 <213> ORGANISM: eucalyptus grandis
169 <220> FEATURE:
170 <221> NAME/KEY: misc_feature
171 <222> LOCATION: (8)..(8)
172 <223> OTHER INFORMATION: n represents inosine
175 <220> FEATURE:
176 <221> NAME/KEY: misc_feature
177 <222> LOCATION: (5)..(5)
178 <223> OTHER INFORMATION: n represents inosine
181 <220> FEATURE:
182 <221> NAME/KEY: misc_feature
183 <222> LOCATION: (11)..(11)
184 <223> OTHER INFORMATION: n represents a, g, t, or c
187 <220> FEATURE:
188 <221> NAME/KEY: misc_feature
189 <222> LOCATION: (17)..(17)
190 <223> OTHER INFORMATION: n represents a, g, t, or c
193 <400> SEQUENCE: 7
W--> 194 atggnggngc ntgtggnta 19
197 <210> SEQ ID NO: 8
198 <211> LENGTH: 20
199 <212> TYPE: DNA
200 <213> ORGANISM: eucalyptus grandis
202 <220> FEATURE:
203 <221> NAME/KEY: misc_feature
204 <222> LOCATION: (12)..(12)
205 <223> OTHER INFORMATION: n = a, g, t, or c
208 <220> FEATURE:
209 <221> NAME/KEY: misc_feature
210 <222> LOCATION: (9)..(9)
211 <223> OTHER INFORMATION: y = t or c
214 <220> FEATURE:
215 <221> NAME/KEY: misc_feature
216 <222> LOCATION: (6)..(6)
217 <223> OTHER INFORMATION: r = a or g
220 <220> FEATURE:
221 <221> NAME/KEY: misc_feature
222 <222> LOCATION: (18)..(18)
223 <223> OTHER INFORMATION: r = a or g
226 <400> SEQUENCE: 8
W--> 227 tgccarttyt gncccartt 20

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/11/2002
PATENT APPLICATION: US/09/383,579B TIME: 10:38:42

Input Set : A:\2531-1-001revised.ST25.txt
Output Set: N:\CRF4\09112002\I383579B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:7; N Pos. 5,8,11,17
Seq#:8; N Pos. 12

VERIFICATION SUMMARY

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L:194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:227 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0